

AMENDMENT

Please amend the application without prejudice, without admission, without surrender of subject matter and without intention of creating any estoppel as to equivalents, as follows.

In the Claims

1. (Currently amended) ~~A binding moiety comprising at least one modified monomeric non-antibody ligand V-like domain (VLD) derived from a non-antibody ligand, the at least one monomeric V-like domain being characterised in that comprising~~ at least one CDR loop structure or part thereof that is modified or replaced, such that the solubility of the modified VLD is improved, when compared with an the unmodified VLD, wherein the monomeric VLD is a binding moiety.

2. (Currently amended) ~~A binding moiety~~ The modified VLD according to claim 1, wherein the in which at least one CDR loop structure or part thereof is modified or replaced such that

- (i) the size of the CDR loop structure is increased when compared with the corresponding CDR loop structure in the unmodified VLD; and/or
- (ii) the modification or replacement results in the formation of a disulphide bond within or between one or more of the CDR loop structures.

3. (Currently amended) ~~A binding moiety comprising at least one modified monomeric non-antibody ligand V-like domain (VLD) derived from a non-antibody ligand, the at least one monomeric V-like domain being characterised in that comprising~~ at least one CDR loop structure or part thereof that is modified or replaced, such that

- (i) the size of the CDR loop structure is altered when compared with the corresponding CDR loop structure in an the unmodified VLD; and/or
- (ii) the modification or replacement results in the formation of a disulphide bond within or between one or more of the CDR loop structures.

4. (Currently amended) ~~A binding moiety~~ The modified VLD according to claim 3, wherein in which the size of the CDR loop structure is increased by at least two amino acid residues.

5. (Currently amended) ~~A binding moiety~~ The modified VLD according to claim 3, wherein in which the size of the CDR loop structure is increased by at least six amino acid residues.